## **RAW SEQUENCE LISTING**

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

10/562, 106
TEWP.
01/04/2006

## ENTERED



**IFWP** 

RAW SEQUENCE LISTING DATE: 01/04/2006
PATENT APPLICATION: US/10/562,106 TIME: 10:24:51

Input Set : A:\466986.txt

Output Set: N:\CRF4\01042006\J562106.raw

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3 <110> APPLICANT: Agrinomics LLC
     5 <120> TITLE OF INVENTION: GENERATION OF PLANTS WITH IMPROVED DROUGHT TOLERANCE
     7 <130> FILE REFERENCE: 6616-72617-02
C--> 9 <140> CURRENT APPLICATION NUMBER: US/10/562,106
     9 <141> CURRENT FILING DATE: 2005-12-22
     9 <150> PRIOR APPLICATION NUMBER: 60/482,139
    10 <151> PRIOR FILING DATE: 2003-06-24
    12 <150> PRIOR APPLICATION NUMBER: PCT/US04/20321
    13 <151> PRIOR FILING DATE: 2004-06-23
    15 <160> NUMBER OF SEQ ID NOS: 3
    17 <170> SOFTWARE: PatentIn version 3.2
    19 <210> SEQ ID NO: 1
    20 <211> LENGTH: 600
    21 <212> TYPE: DNA
    22 <213> ORGANISM: Arabidopsis thaliana
    24 <400> SEQUENCE: 1
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                                                                              120
    29 ttcatcaacc acagaggatc cgacacaaag agaaacatcg caacattgct ttatgacaat
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    31 ctcaaatctc gtaacttacg tccattcttg gattccaaga acatgaagcc cggagataag
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    33 ctttttgatc atatcaacaa cgcgattctc acttctaaag tcgctgtaac ggttttttct
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    35 cctaactact gcgattcata tttctgtctg cacgagcttg ctcttataat ggagtccaga
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    37 aaaagggtca taccgatatt ctgcgacatc aaaccttcac aactcgatgt tatgatcgag
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    39 agggtgacat gttctgatga tgaaatccaa aggtttagat gggcccttca agaagctaaa
                                                                              480
    41 gatategttg gacteaegtt tgatteetae aaagggaatt tateggaggt tgttaeagtt
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    43 gcatcggatg ttatcgtcga gagattggtc gagttagagg ctaaagatga aaatctatag
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    47 <211> LENGTH: 199
    48 <212> TYPE: PRT
    49 <213> ORGANISM: Arabidopsis thaliana
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    53 Met Ile Ser Arg Val Phe Asn Phe Met Lys Gln Ser Glu Thr Val Gln
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    54 1
                                                                 15
                        5
    57 Leu Asn Asn Lys Leu Phe Leu Asp Leu Leu Ser Ser Ser Ser Ala
    58
                   20
                                        25
    61 Lys Pro Lys Ile Leu His Asp Val Phe Ile Asn His Arg Gly Ser Asp
               35
                                    40
    65 Thr Lys Arg Asn Ile Ala Thr Leu Leu Tyr Asp Asn Leu Lys Ser Arg
                                55
    69 Asn Leu Arg Pro Phe Leu Asp Ser Lys Asn Met Lys Pro Gly Asp Lys
                            70
                                                75
    73 Leu Phe Asp His Ile Asn Asn Ala Ile Leu Thr Ser Lys Val Ala Val
    74
                        85
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129 gaggaggtta agttcaccgt gggactcacg ttcaactcct caaaa

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77 Thr Val Phe Ser Pro Asn Tyr Cys Asp Ser Tyr Phe Cys Leu His Glu
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81 Leu Ala Leu Ile Met Glu Ser Arg Lys Arg Val Ile Pro Ile Phe Cys
82
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                               120
                                                    125
85 Asp Ile Lys Pro Ser Gln Leu Asp Val Met Ile Glu Arg Val Thr Cys
                           135
                                                140
89 Ser Asp Asp Glu Ile Gln Arg Phe Arg Trp Ala Leu Gln Glu Ala Lys
                       150
                                           155
93 Asp Ile Val Gly Leu Thr Phe Asp Ser Tyr Lys Gly Asn Leu Ser Glu
                   165
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97 Val Val Thr Val Ala Ser Asp Val Ile Val Glu Arg Leu Val Glu Leu
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101 Glu Ala Lys Asp Glu Asn Leu
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106 <211> LENGTH: 585
107 <212> TYPE: DNA
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111 ttaatcatca agaatacact cacatacata catacataga acacctctat tgttataaat
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113 tectagetat atattttett aageageeae accatgeaae gtaacatgge aatgagette
                                                                          120
115 ttccagcgca ggttcttcag ccagcaaagg cgaacacaaa tggtggcaag gcgagtgttg
                                                                          180
117 gagccgtgtg acgtgttcct gaaccatagg agcatggaca caaagaaaac agtagccacg
                                                                          240
119 ttgctctatg accatttgaa gaggcacggt ttcaaccctt tcttggacaa caagaacatg
                                                                          300
121 aagccagggg acaaactgtt tgagaaaatc aatagggctg ttatggagtg caagattggg
                                                                          360
123 gtggcggttc tctcgccacg ctacacggag tcctattttt gcctccatga gcttgcactt
                                                                          420
125 cttctggggt gcaacaagaa ggtcattccc atcttctgtg atgtaaagcc ttcgcagttg
                                                                          480
127 egegttgtca acaaccetaa gtggtetgaa gatgaactta gacgetteag acgggetett
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585

VERIFICATION SUMMARY

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L:9 M:270 C: Current Application Number differs, Replaced Current Application No

L:9 M:271 C: Current Filing Date differs, Replaced Current Filing Date